Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in this application:

- 1. (currently amended) An isolated complex comprising: a An isolated bacterial heme binding protein complexed with a porphyrin, wherein said protein complex reversibly binds oxygen with a low affinity and wherein said protein comprises a heme binding domain that associates with the porphyrin and an aerotaxis signaling domain of said protein that shows at least 20% identity to a myoglobin heme binding domain having an amino acid sequence of SEQ ID NO:76.
 - 2-5 (canceled)
- 6. (currently amended) The isolated heme binding protein complex according to claim 1, wherein the protein has an amino acid sequence of SEQ ID NO:2.
 - 7-10 (canceled)
- 11. (currently amended) A blood substitute <u>comprising</u> eomprising: a <u>complex according to claim 1</u> bacterial heme binding protein wherein said protein reversibly binds oxygen with a low affinity and comprises a heme binding domain that shows at least 20% identity to a myoglobin heme binding domain having an amino acid sequence of SEQ ID NO:76.
 - 12-15 (canceled)
- 16. (previously presented) The blood substitute according to claim 11, wherein the protein has an amino acid sequence of SEQ ID NO:2.
 - 17-47 (canceled)
 - 48. (original) A chimeric protein comprising: a heme-binding domain of an isolated heme binding bacterial protein; and a heterologous signaling domain.
- 49. (previously presented) The chimeric protein according to claim 48, wherein the heterologous signaling domain is a mutated signaling domain having altered affinity for its ligand.

- 50. (canceled)
- 51. (previously presented) The chimeric protein according to claim 48, wherein the heme binding domain is from a heme binding protein isolated from *Archaea*.
- 52. (previously presented) The chimeric protein according to claim 51, wherein the heme binding protein is isolated from *Halobacterium salinarium*.
- 53. (previously presented) The chimeric protein according to claim 52, wherein the activity of the heme binding protein is salt tolerant.
- 54. (currently amended) The chimeric protein according to claim <u>52</u> 53, wherein the protein has an <u>the heme binding domain comprises the</u> amino acid sequence of <u>SEQ ID NO:2 SEQ ID NO: 77</u>.
 - 55-65 (canceled)
- 66. (currently amended) The <u>complex</u> isolated heme binding protein according to claim 1 wherein the <u>complex</u> protein is purified.
- 67. (currently amended) The <u>complex</u> isolated heme binding protein according to claim 1 wherein the <u>complex</u> protein is recombinant.
- 68. (new) The complex according to claim 1, wherein the heme binding domain comprises a plurality of α -helices.
- 69. (new) The complex according to claim 68, wherein the heme binding domain comprises eight α -helices.
- 70. (new) The complex according to claim 1, wherein the heme binding domain is positioned N-terminal and the aerotaxis signaling domain is positioned C-terminal in the heme binding protein.
- 71. (new) The complex according to claim 1, wherein the heme binding domain is at least 20% identical to SEQ ID NO: 76.

- 72. (new) The complex according to claim 1, wherein the aerotaxis signaling domain is at least 30% identical to SEQ ID NO: 79.
- 73. (new) The complex according to claim 1, wherein said protein is about 50 kDa.
- 74. (new) The complex according to claim 1, wherein the porphyrin is a Fe-porphyrin.
- 75. (new) The complex according to claim 74, wherein the Fe-porphyrin is a heme molecule.
- 76. (new) The complex according to claim 75, wherein the heme molecule is a b-type heme molecule.
- 77. (new) The complex according to claim 75, wherein the complex has an oxygenated form characterized as having spectral properties of: Soret band absorption at 406 nm, α -band absorption at 578 nm, and β -band absorption at 538 nm.
- 78. (new) The complex according to claim 75, wherein the complex has a deoxygenated form characterized as having spectral properties of: Soret band absorption at 425 nm, and converged α -band and β band absorption centered at 555 nm.
- 79. (new) The complex according to claim 1, wherein the porphyrin is a Zn-porphyrin.
- 80. (new) The complex according to claim 1, wherein the porphyrin is a Sn-porphyrin.
- 81. (new) The blood substitute according to claim 11, wherein the porphyrin is a Fe-porphyrin.
- 82. (new) The blood substitute according to claim 81, wherein the Feporphyrin is a heme molecule.